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Timestamp: [year=2008; month=4; day=29; hr=12; min=4; sec=59; ms=585;]

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Application No: 10672399 Version No: 2.0

Input Set:

Output Set:

Started: 2008-04-15 15:40:46.474

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Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 445 ms

Total Warnings: 0

Total Errors: 0

No. of SeqIDs Defined: 15

Actual SeqID Count: 15

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<120> Cancer Monitoring and Therapeutics

<130> 05-013

<140> 10672399

<141> 2003-09-25

<150> 60/472,401

<151> 2003-05-22

<160> 15

<170> PatentIn version 3.3

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<211> 1737

<212> DNA

<213> Homo sapiens

<400> 1

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 <211> 578
 <212> PRT
 <213> Homo sapiens

 <400> 2

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Leu Gly Leu Met Thr Trp Ala Tyr Ala Ala Gly Val Pro Leu Ala Ser
 35 40 45

Asp Arg Tyr Gly Leu Leu Ala Phe Gly Leu Tyr Gly Ala Phe Leu Ser
 50 55 60

Ala His Leu Val Ala Gln Ser Leu Phe Ala Tyr Leu Glu His Arg Arg
 65 70 75 80

Val Ala Ala Ala Ala Arg Gly Pro Leu Asp Ala Ala Thr Ala Arg Ser
 85 90 95

Val	Ala	Leu	Thr	Ile	Ser	Ala	Tyr	Gln	Glu	Asp	Pro	Ala	Tyr	Leu	Arg	100	105	110
Gln	Cys	Leu	Ala	Ser	Ala	Arg	Ala	Leu	Leu	Tyr	Pro	Arg	Ala	Arg	Leu	115	120	125
Arg	Val	Leu	Met	Val	Val	Asp	Gly	Asn	Arg	Ala	Glu	Asp	Leu	Tyr	Met	130	135	140
Val	Asp	Met	Phe	Arg	Glu	Val	Phe	Ala	Asp	Glu	Asp	Pro	Ala	Thr	Tyr	145	150	155
Val	Trp	Asp	Gly	Asn	Tyr	His	Gln	Pro	Trp	Glu	Pro	Ala	Ala	Ala	Gly	165	170	175
Ala	Val	Gly	Ala	Gly	Ala	Tyr	Arg	Glu	Val	Glu	Ala	Glu	Asp	Pro	Gly	180	185	190
Arg	Leu	Ala	Val	Glu	Ala	Leu	Val	Arg	Thr	Arg	Arg	Cys	Val	Cys	Val	195	200	205
Ala	Gln	Arg	Trp	Gly	Gly	Lys	Arg	Glu	Val	Met	Tyr	Thr	Ala	Phe	Lys	210	215	220
Ala	Leu	Gly	Asp	Ser	Val	Asp	Tyr	Val	Gln	Val	Cys	Asp	Ser	Asp	Thr	225	230	235
Arg	Leu	Asp	Pro	Met	Ala	Leu	Leu	Glu	Leu	Val	Arg	Val	Leu	Asp	Glu	245	250	255
Asp	Pro	Arg	Val	Gly	Ala	Val	Gly	Gly	Asp	Val	Arg	Ile	Leu	Asn	Pro	260	265	270
Leu	Asp	Ser	Trp	Val	Ser	Phe	Leu	Ser	Ser	Leu	Arg	Tyr	Trp	Val	Ala	275	280	285
Phe	Asn	Val	Glu	Arg	Ala	Cys	Gln	Ser	Tyr	Phe	His	Cys	Val	Ser	Cys	290	295	300
Ile	Ser	Gly	Pro	Leu	Gly	Leu	Tyr	Arg	Asn	Asn	Leu	Leu	Gln	Gln	Phe	305	310	315

Leu Glu Ala Trp Tyr Asn Gln Lys Phe Leu Gly Thr His Cys Thr Phe
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Gly Asp Asp Arg His Leu Thr Asn Arg Met Leu Ser Met Gly Tyr Ala
340 345 350

Thr Lys Tyr Thr Ser Arg Ser Arg Cys Tyr Ser Glu Thr Pro Ser Ser
355 360 365

Phe Leu Arg Trp Leu Ser Gln Gln Thr Arg Trp Ser Lys Ser Tyr Phe
370 375 380

Arg Glu Trp Leu Tyr Asn Ala Leu Trp Trp His Arg His His Ala Trp
385 390 395 400

Met Thr Tyr Glu Ala Val Val Ser Gly Leu Phe Pro Phe Phe Val Ala
405 410 415

Ala Thr Val Leu Arg Leu Phe Tyr Ala Gly Arg Pro Trp Ala Leu Leu
420 425 430

Trp Val Leu Leu Cys Val Gln Gly Val Ala Leu Ala Lys Ala Ala Phe
435 440 445

Ala Ala Trp Leu Arg Gly Cys Leu Arg Met Val Leu Leu Ser Leu Tyr
450 455 460

Ala Pro Leu Tyr Met Cys Gly Leu Leu Pro Ala Lys Phe Leu Ala Leu
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Val Thr Met Asn Gln Ser Gly Trp Gly Thr Ser Gly Arg Arg Lys Leu
485 490 495

Ala Ala Asn Tyr Val Pro Leu Leu Pro Leu Ala Leu Trp Ala Leu Leu
500 505 510

Leu Leu Gly Gly Leu Val Arg Ser Val Ala His Glu Ala Arg Ala Asp
515 520 525

Trp Ser Gly Pro Ser Arg Ala Ala Glu Ala Tyr His Leu Ala Ala Gly
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Ala Gly Ala Tyr Val Gly Tyr Trp Val Ala Met Leu Thr Leu Tyr Trp

545 550 555 560

Val Gly Val Arg Arg Leu Cys Arg Arg Arg Thr Gly Gly Tyr Arg Val
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Gln Val

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<211> 1071
<212> DNA
<213> Homo sapiens

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<210> 4
<211> 376
<212> PRT

<213> Homo sapiens

<400> 4

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Leu Gly Leu Met Thr Trp Ala Tyr Ala Ala Gly Val Pro Leu Ala Ser
35 40 45

Asp Arg Tyr Gly Leu Leu Ala Phe Gly Leu Tyr Gly Ala Phe Leu Ser
50 55 60

Ala His Leu Val Ala Gln Ser Leu Phe Ala Tyr Leu Glu His Arg Arg
65 70 75 80

Val Ala Ala Ala Ala Arg Gly Pro Leu Asp Ala Ala Thr Ala Arg Ser
85 90 95

Val Ala Leu Thr Ile Ser Ala Tyr Gln Glu Asp Pro Ala Tyr Leu Arg
100 105 110

Gln Cys Leu Ala Ser Ala Arg Ala Leu Leu Tyr Pro Arg Ala Arg Leu
115 120 125

Arg Val Leu Met Val Val Asp Gly Asn Arg Ala Glu Asp Leu Tyr Met
130 135 140

Val Asp Met Phe Arg Glu Val Phe Ala Asp Glu Asp Pro Ala Thr Tyr
145 150 155 160

Val Trp Asp Gly Asn Tyr His Gln Pro Trp Glu Pro Ala Ala Ala Gly
165 170 175

Ala Val Gly Ala Gly Ala Tyr Arg Glu Val Glu Ala Glu Asp Pro Gly
180 185 190

Arg Leu Ala Val Glu Ala Leu Val Arg Thr Arg Arg Cys Val Cys Val
195 200 205

Ala Gln Arg Trp Gly Gly Lys Arg Glu Val Met Tyr Thr Ala Phe Lys

210

215

220

Ala Leu Gly Asp Ser Val Asp Tyr Val Gln Val Cys Asp Ser Asp Thr
 225 230 235 240

Arg Leu Asp Pro Met Ala Leu Leu Glu Leu Val Arg Val Leu Asp Glu
 245 250 255

Asp Pro Arg Val Gly Ala Val Gly Gly Asp Val Arg Ile Leu Asn Pro
 260 265 270

Leu Asp Ser Trp Val Ser Phe Leu Ser Ser Leu Arg Tyr Trp Val Ala
 275 280 285

Phe Asn Val Glu Arg Ala Cys Gln Ser Tyr Phe His Cys Val Ser Cys
 290 295 300

Ile Ser Gly Ser Leu Gly Thr Pro Pro Gly Pro Ala Ala Thr Gln Arg
 305 310 315 320

Arg Pro Arg Pro Ser Cys Gly Gly Ala Ser Arg His Ala Gly Pro Ser
 325 330 335

Arg Thr Ser Val Ser Gly Cys Thr Thr Arg Ser Gly Gly Thr Gly Thr
 340 345 350

Met Arg Gly Pro Thr Arg Arg Trp Ser Pro Ala Cys Ser Pro Ser Ser
 355 360 365

Trp Arg Pro Leu Cys Cys Val Cys
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<210> 5

<211> 1080

<212> DNA

<213> Homo sapiens

<400> 5

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gccttccttt cagcgcacct ggtggcgcag agcctcttcg cgtacctgga gcaccggcgg 240

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<210> 6
 <211> 360
 <212> PRT
 <213> Homo sapiens

<400> 6

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Leu Gly Leu Met Thr Trp Ala Tyr Ala Ala Gly Val Pro Leu Ala Ser
 35 40 45

Asp Arg Tyr Gly Leu Leu Ala Phe Gly Leu Tyr Gly Ala Phe Leu Ser
 50 55 60

Ala His Leu Val Ala Gln Ser Leu Phe Ala Tyr Leu Glu His Arg Arg
 65 70 75 80

Val Ala Ala Ala Ala Arg Gly Pro Leu Asp Ala Ala Thr Ala Arg Ser

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90

95

Val Ala Leu Thr Ile Ser Ala Tyr Gln Glu Asp Pro Ala Tyr Leu Arg
100 105 110

Gln Cys Leu Ala Ser Ala Arg Ala Leu Leu Tyr Pro Arg Ala Arg Leu
115 120 125

Arg Val Leu Met Val Val Asp Gly Asn Arg Ala Glu Asp Leu Tyr Met
130 135 140

Val Asp Met Phe Arg Glu Val Phe Ala Asp Glu Asp Pro Ala Thr Tyr
145 150 155 160

Val Trp Asp Gly Asn Tyr His Gln Pro Trp Glu Pro Ala Ala Ala Gly
165 170 175

Ala Val Gly Ala Gly Ala Tyr Arg Glu Val Glu Ala Glu Asp Pro Gly
180 185 190

Arg Leu Ala Val Glu Ala Leu Val Arg Thr Arg Arg Cys Val Cys Val
195 200 205

Ala Gln Arg Trp Gly Gly Lys Arg Glu Val Met Tyr Thr Ala Phe Lys
210 215 220

Ala Leu Gly Asp Ser Val Asp Tyr Val Gln Val Cys Asp Ser Asp Thr
225 230 235 240

Arg Leu Asp Pro Met Ala Leu Leu Glu Leu Val Arg Val Leu Asp Glu
245 250 255

Asp Pro Arg Val Gly Ala Val Gly Gly Asp Val Arg Ile Leu Asn Pro
260 265 270

Leu Asp Ser Trp Val Ser Phe Leu Ser Ser Leu Arg Tyr Trp Val Ala
275 280 285

Phe Asn Val Glu Arg Ala Cys Gln Ser Tyr Phe His Cys Val Ser Cys
290 295 300

Ile Ser Gly Pro Leu Glu Ser Cys Pro Gly Pro Arg Glu His Ala Met
305 310 315 320

Met Pro Ser Phe Leu Ala Pro Val Gln Val His Leu Gln Val Pro Leu
325 330 335

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340 345 350

Thr Leu Val Gln Val Val Leu Pro
355 360

<210> 7
<211> 1065
<212> DNA
<213> Homo sapiens

<400> 7
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